



532

99/787323
PCT/PTO 16-MAR 2001**ABSTRACT OF THE DISCLOSURE**

5

Multiple Sequencing Method

a' 7

10 The present invention provides a method for identifying a nucleic acid utilizing a run-off sequencing reaction of a relatively short portion of the nucleic acid. The method can be utilized, for example, to identify an EST from only a small portion of the EST and in an analysis of nucleotide polymorphisms.

09/03/2001 10:00:00